

Raw Sequence Listing before editing,
for reference only

BEST AVAILABLE COPY



PCT

RAW SEQUENCE LISTING

DATE: 05/12/2005

PATENT APPLICATION: US/10/533,750

TIME: 08:04:08

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\05122005\J533750.raw

3 <110> APPLICANT: Yoichi Matsubara
 4 Shigeo Kure
 5 Mitsubishi Kagaku Bio-Clinical Laboratories, Inc.
 7 <120> TITLE OF INVENTION: Method of detecting gene mutation
 W--> 0 <130> FILE REFERENCE:
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/533,750
 C--> 10 <141> CURRENT FILING DATE: 2005-05-04
 12 <150> PRIOR APPLICATION NUMBER: JP 2002-323419
 13 <151> PRIOR FILING DATE: 2002-11-07
 15 <160> NUMBER OF SEQ ID NOS: 64

ERRORED SEQUENCES

711 <210> SEQ ID NO: 64
 712 <211> LENGTH: 12
 713 <212> TYPE: DNA
 714 <213> ORGANISM: Artificial Sequence
 716 <220> FEATURE:
 717 <223> OTHER INFORMATION: probe
 719 <400> SEQUENCE: 64
 720 acaggcgagg aa
 E--> 726 1/16
 E--> 727 118308.1

Does Not Comply
Corrected Diskette Needed

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VERIFICATION SUMMARY

DATE: 05/12/2005

PATENT APPLICATION: US/10/533,750

TIME: 08:04:09

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\05122005\J533750.raw

L:0 M:201 W: Mandatory field data missing, <130> FILE REFERENCE
L:9 M:270 C: Current Application Number differs, Replaced Current Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:726 M:254 E: No. of Bases conflict, LENGTH:Input:16 Counted:13 SEQ:64
L:726 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
M:254 Repeated in SeqNo=64
L:727 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:7
L:727 M:252 E: No. of Seq. differs, <211> LENGTH:Input:12 Found:14 SEQ:64